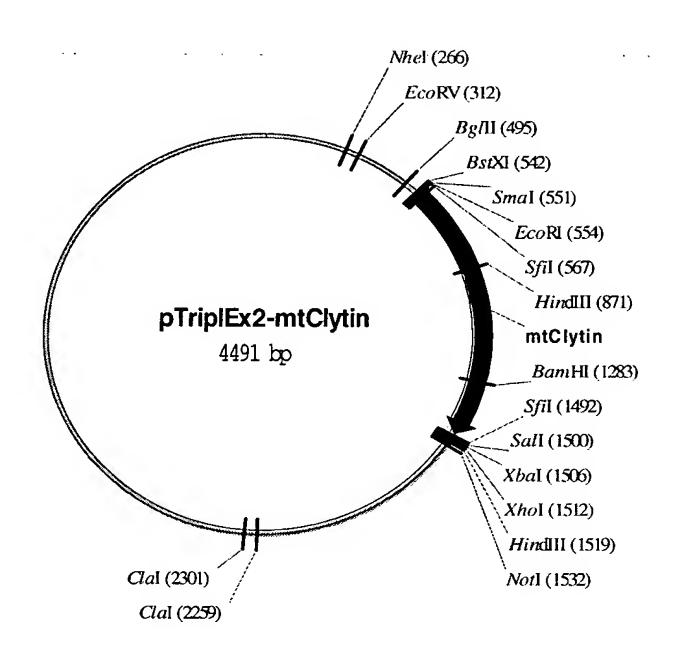
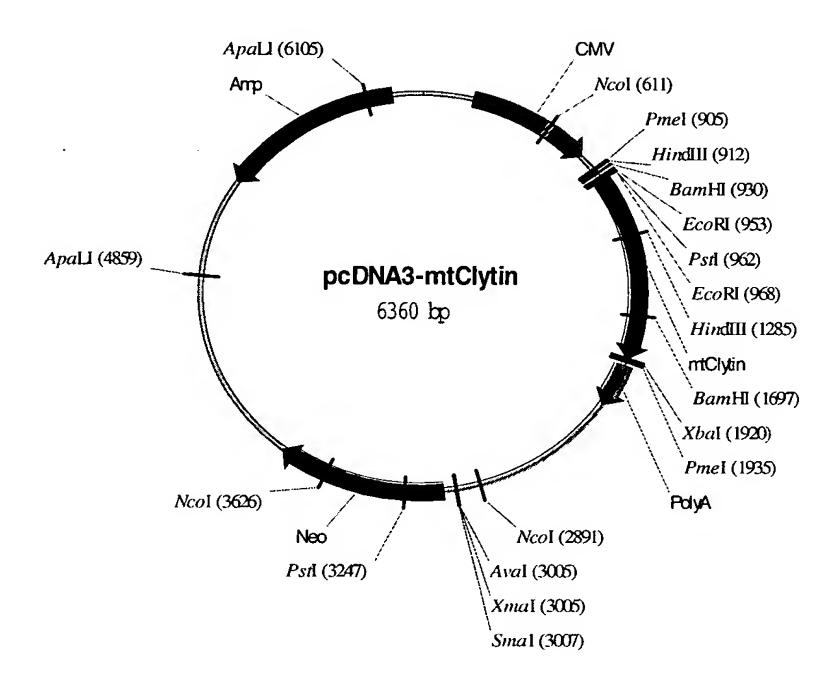
## **Figures**

Fig. 1



<u>Fig. 2</u>



<u>Fig. 3</u>

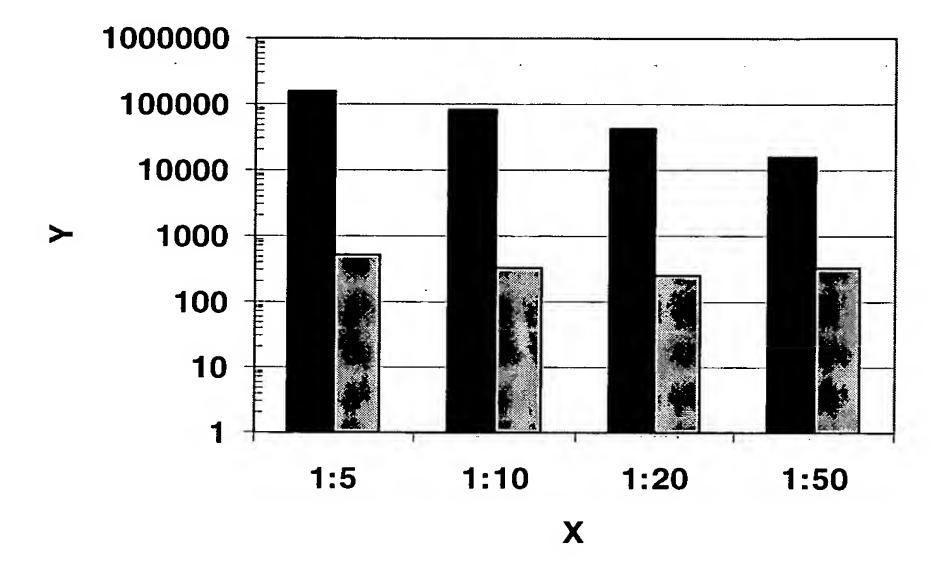


Fig. 4

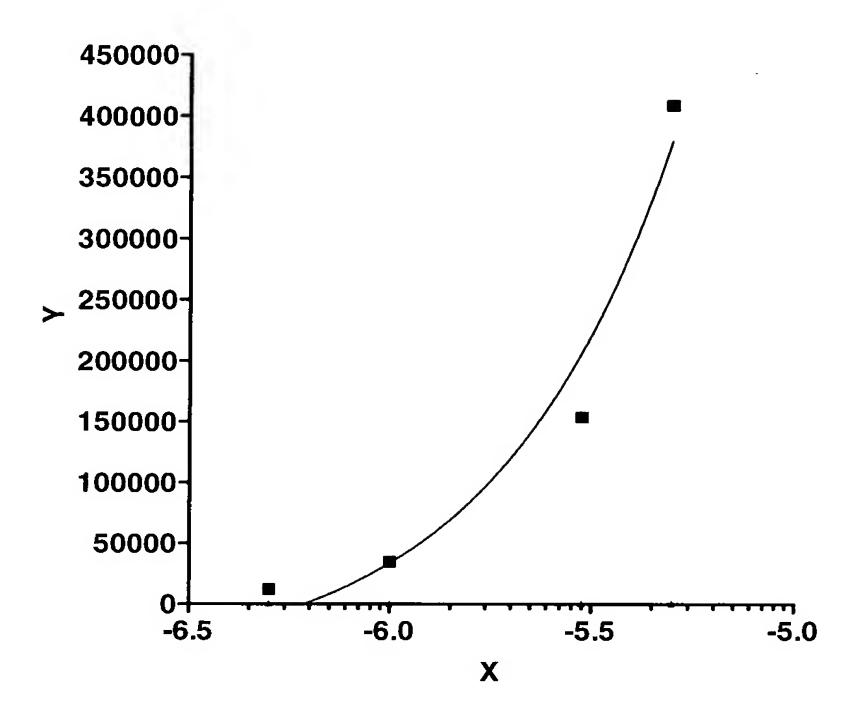


Fig. 5

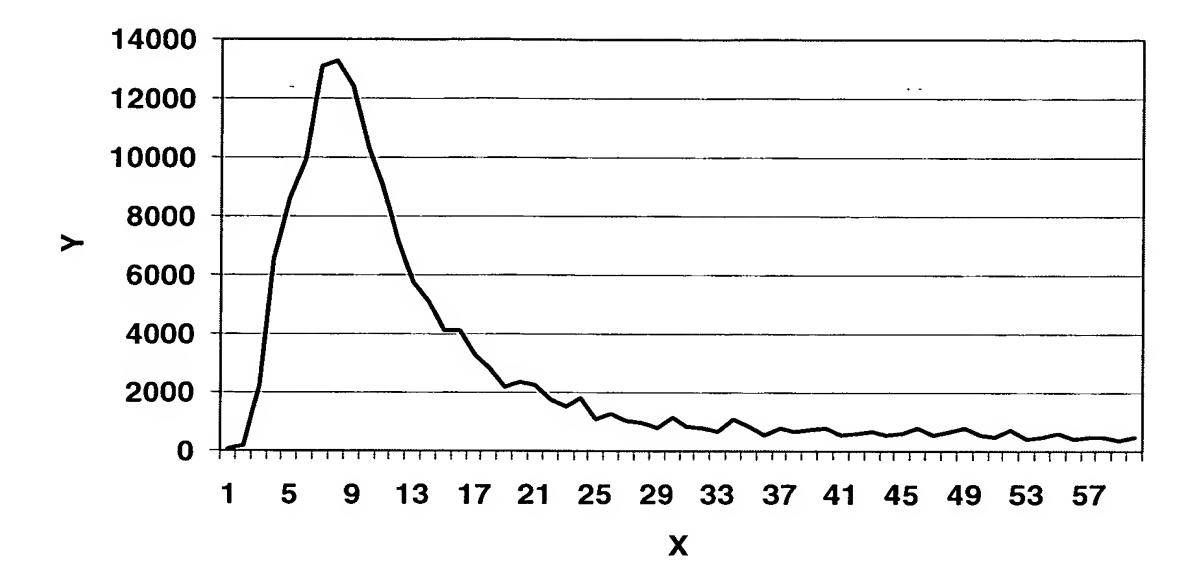
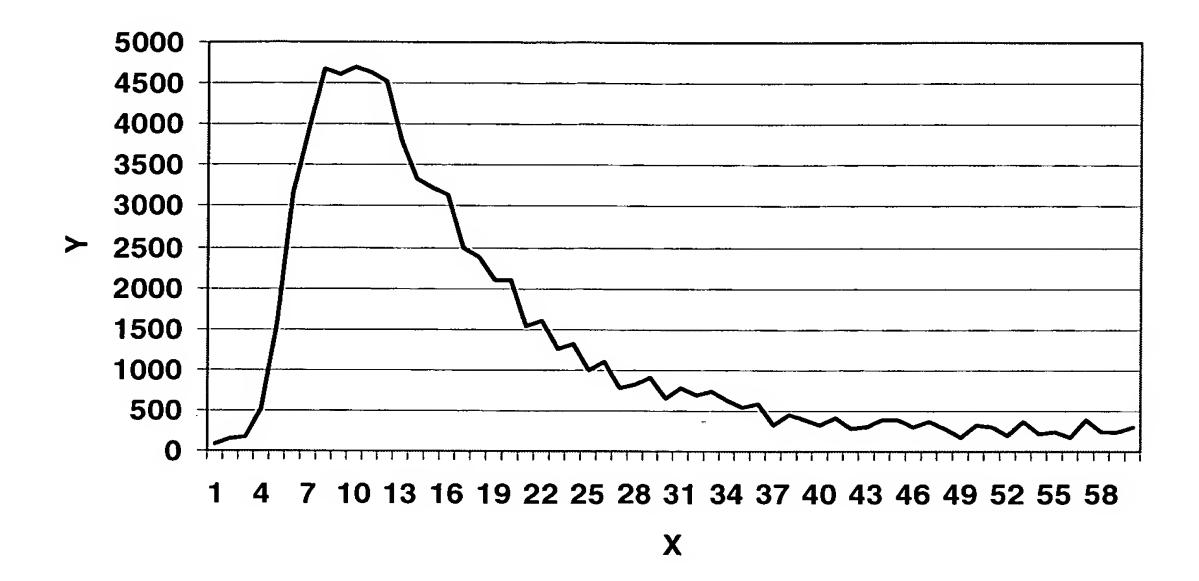


Fig. 6



## Fig. 7

1				50	
Clytin					• • • • • • • • • • • • • • • • • • • •
mtClytin	GACAGATAAA	AAATTCACTC	CTTAGATTAT	' TTAGTGAATA	AGAGAAAA
	51				100
Clytin	• • • • • • • • • • • • • • • • • • • •	• • • • • • • • • •			
mtClytin	AGGATAAGAA	ATCAAGATGC	AAAGGTTTAC	AAATCGTCTT	CTTTCCATG
	101				150
Clytin		ATCA	ACTTTTGCAA	CTCAAAGCAA	ATTTCAAAAC
mtClytin	CGGCTTTACG	TGCAAGATCA	AGATT.GCAA	CGCACGGCAA	ATTTTCACAC
	151				200
Clytin	TTCAACATGG	CTGAC.ACTG	САТСААААТА	CGCCGTCAAA	CTCAGACCC
mtClytin	CAGCATACTC	TTGGCTACAG	ATTCAAAATA	CGCGGTCAAA	CTCGATCCTC
	201				250
Clytin	ACTTCGACAA	CCCAAAATGG	GTCAACAGAC	ACAAATTTAT	GTTCAACTTI
mtClytin	ATTTTGCAAA	TCCAAAATGG	ATCAACAGAC	ACAAATTTAT	GTTCAACTTT
	251				300
Clytin	TTGGACATTA	ACGGCGACGG	AAAAATCACT	TTGGATGAAA	TCGTCTCCAA
mtClytin	TTGGACATAA	ACGGTAAGGG	GAAAATCACA	TTAGATGAAA	TCGTCTCCAA
	301				350
Clytin	AGCTTCGGAT	GACATTTGCG	CCAAACTTGG	AGCAACACCA	GAACAGACCA
mtClytin	AGCTTCAGAC	GACATTTGTG	CTAAACTGGA	TGCAACACCA	GAACAGACCA
	351				400
Clytin	AACGTCACCA	GGATGCTGTC	GAAGCTTTCT	TCAAAAAGAT	TGGTATGGAT
mtClytin	AACGTCACCA	GGATGCTGTT	GAAGCCTTTT	TCAAGAAAAT	GGGCATGGAT
	401				450
Clytin	TATGGTAAAG	AAGTCGAATT	CCCAGCTTTT	GTTGATGGAT	GGAAAGAACT
mtClytin	TATGGTAAAG	AAGTTGCATT	CCCAGAATTT	ATTAAGGGAT	GGGAAGAGTT
	451				500
Clytin	GGCCAATTAT	GACTTGAAAC	TTTGGTCTCA	AAACAAGAAA	TCTTTGATCC
mtClytin	GGCCGAACAC	GACTTGGAAC	TCTGGTCTCA	AAACAAAAGT	ACATTGATCC
	501				550
Clytin	GCGACTGGGG	AGAAGCTGTT	TTCGACATTT	TTGACAAAGA	CGGAAGTGGC
mtClytin	GTGAATGGGG	AGATGCTGTT	TTCGACATTT	TCGACAAAGA	CGCAAGTGGC

	551				600
Clytin	TCAATCAGTT	TGGACGAATG	GAAGGCTTAT	GGACGAATCT	CTGGAATCTG
mtClytin	TCAATCAGTT	TAGACGAATG	GAAGGCTTAC	GGACGAATCT	CTGGAATCTG
	601				650
Clytin	CTCATCAGAC	GAAGACGCCG	AAAAGACCTT	CAAACATTGC	GATTTGGACA
mtClytin	TCCATCAGAC	GAAGACGCTG	AGAAGACGTT	CAAACATTGT	GATTTGGACA
	651				700
Clytin	ACAGTGGCAA	ACTTGATGTT	GATGAGATGA	CCAGACAACA	TTTGGGATTC
mtClytin	ACAGTGGCAA	ACTTGATGTT	GATGAGATGA	CCAGGCAACA	TTTAGGCTTC
	701				750
Clytin	TGGTACACCT	TGGACCCCAA	CGCTGATGGT	CTTTACGGCA	ATTTTGTTCC
mtClytin	TGGTACACAT	TGGATCCAAC	TTCTGATGGT	CTTTATGGCA	ATTTTGTTCC
	751				800
Clytin				AAGTTTTGGA	
mtClytin	CTAAGAAGCG	TTCAGTTAAA	AACGCTAAAC	ATTGTTCAGT	TGTAAAATTA
	0.01				
<b>61</b> 4.1	801				850
	GATACTAT				
mtClytin	TATTCATTTT	CATTTCGTAA	AATTAGTATT	TATAAATTTG	TATCATAAAT
	851				000
Clutin			3 3 CO 3 3 COCOCOCO		900
	TGTAAC.ATG				
mcciytin	TGTATCCATG	TIGTAGACTA	AATAAGACTC	GGCAAAAAA	AAAAAAAAA
	901	913			
Clytin					
	AAAAAAAA				
·	·	- <del></del>			

Fig. 8			
	1	50	
mtClytin	MQRFTNRLLS MSALRARSRL QRTANFH	TSI LLATDSKYAV KLDPDFANPK	
Clytin	••••••	MADTASKYAV KLRPNFDNPK	
	51	100	
mtCyltin	WINRHKFMFN FLDINGKGKI TLDEIVS	KAS DDICAKLDAT PEQTKRHQDA	
Clytin	WVNRHKFMFN FLDINGDGKI TLDEIVS	KAS DDICAKLGAT PEQTKRHQDA	
	101	150	
Clytin	VEAFFKKMGM DYGKEVAFPE FIKGWEE		
Clytin	VEAFFKKIGM DYGKEVEFPA FVDGWKE	LAN YDLKLWSQNK KSLIRDWGEA	
	151	200	
Clytin	VFDIFDKDAS GSISLDEWKA YGRISGIO	CPS DEDAEKTFKH CDLDNSGKLD	
Clytin	VFDIFDKDGS GSISLDEWKA YGRISGIO	CSS DEDAEKTFKH CDLDNSGKLD	
	201	228	
mtClytin	-		
Clytin	VDEMTRQHLG FWYTLDPNAD GLYGNFVI	<u>-</u>	
Fig. 9			
1			50
mtClytir	n MQRFTNRLLS MSALRARSRL	QRTANFHTSI LLATDSKYA	V KLDPDFANPK
clytin-2	2	MTDTASKYA	V KLKTNFEDPK
Clytin	n	MADTASKYA	V KLRPNFDNPK
	51		100
mtClytin			
clytin-2			
Clytin	n WVNRHKFMFN FLDINGDGKI	TLDEIVSKAS DDICAKLGA	r PEQTKRHQDA
	101		150
mtClytin	n VEAFFKKMGM DYGKEVAFPE	FIKGWEELAE HDLELWSQN	
clytin-2	2 VEAFFKKIGL DYGKEVEFPA	FVNGWKELAK HDLKLWSQNI	K KSLIRNWGEA
Clytin	n VEAFFKKIGM DYGKEVEFPA	FVDGWKELAN YDLKLWSQNI	K KSLIRDWGEA
	151		200
mtClytin	vFDIFDKDAS GSISLDEWKA	YGRISGICPS DEDAEKTFKI	H CDLDNSGKLD
clytin-2	2 VFDIFDKDGS GSISLDEWKT	YGGISGICPS DEDAEKTFK	H CDLDNSGKLD
Clytin	vFDIFDKDGS GSISLDEWKA	YGRISGICSS DEDAEKTFKI	H CDLDNSGKLD
	201	228	
mtClytin			
clytin-2	~		
Clytin	<del>-</del>		
~ + 7 + + + + + + + + + + + + + + + + +	· ANTITIVATIO LATITURIAN (	ONIGHEAE	